Mal.Oecy−Bulk_difneg domain

CLIP	p = 0.00672 n = 12
MCM	p = 0.00672 n = 7
MCM_OB	p = 0.00672 n = 7
zf−MYND	p = 0.0128 n = 15
DHC_N2	p = 0.0131 n = 8
Trypsin	p = 0.0133 n = 70
AAA_6	p = 0.0145 n = 7
AAA_8	p = 0.0145 n = 7
Kinesin	p = 0.0145 n = 10
SET	p = 0.0145 n = 15
AAA_7	p = 0.0192 n = 7
AAA_9	p = 0.0192 n = 7
Dynein_heavy	p = 0.0192 n = 7
MT	p = 0.0192 n = 7
MutS_III	p = 0.0192 n = 4
MutS_V	p = 0.0192 n = 4
Sugar_tr	p = 0.0296 n = 16
OS−D	p = 0.0376 n = 5
PTCB−BRCT	p = 0.0721 n = 4
ARL2_Bind_BART	p = 0.0823 n = 3
DUF1126	p = 0.0823 n = 3
DUF4749	p = 0.0823 n = 3
ERCC4	p = 0.0823 n = 3
FAD_binding_8	p = 0.0823 n = 3
Ferric_reduct	p = 0.0823 n = 3
KIAA1430	p = 0.0823 n = 3
NAD_binding_6	p = 0.0823 n = 3
Polysacc_deac_1	p = 0.0823 n = 3
DOMON	p = 0.0838 n = 6
CBM_14	p = 0.0967 n = 24

0   −1   −2   −3   −4   −5
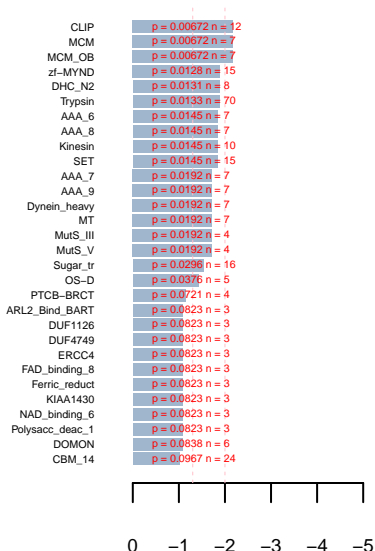
log(p)

n = 1757 genes | p adjust BH TRUE